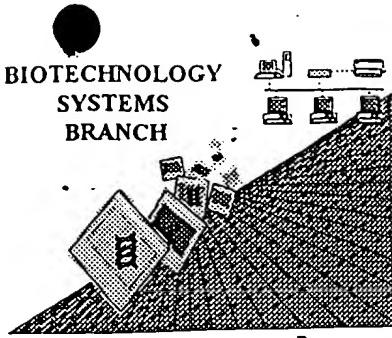


255010390  
316

## RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/754,853

Source: O1PE

Date Processed by STIC: 1/23/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

### Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be downloaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

OIPE

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/754,853

DATE: 01/23/2001  
TIME: 15:39:47

Input Set : D:\pa\_00330.txt  
Output Set: N:\CRF3\01232001\I754853.raw

1 <110> APPLICANT: Parnell, Laurence D.  
2 Hauge, Brian M.  
3 Parsons, Jeremy D.  
4 Wang, Ming Li

6 <120> TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With

7 Soybean Cyst Nematode Resistance  
9 <130> FILE REFERENCE: 38-10(15810)B

*OK* 11 <140> CURRENT APPLICATION NUMBER: US/09/754,853

11 <141> CURRENT FILING DATE: 2001-01-05

11 <150> PRIOR APPLICATION NUMBER: US 60/174,880

13 <151> PRIOR FILING DATE: 2000-01-07

15 <160> NUMBER OF SEQ ID NOS: 1123

17 <210> SEQ ID NO: 1

18 <211> LENGTH: 127197

19 <212> TYPE: DNA

20 <213> ORGANISM: Glycine max

22 <223> OTHER INFORMATION: Seq ID: 515002\_region\_G2

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32 catttttag ttaatttcat caccatcata cgttgtaaat gctatataat taatataatca 240

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36 cactttgtt ttaacatatt atttattatt aattaaaatt tattgaaaac cataaattat 360

38 aaattataag taaaaccgta aaataaagag tcatatcaa attttttgtt gattttcaac 420

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44 taccatctaa tgaagagaga gttttcctaa ctttgcac acagccgtca caatccatca 600

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54 atagagaaaa agcaagggtcc aaaaccacaa gaagttacaa ggaactttct t gcaaaacag 900

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82 gataaaaattt cagtcaactt aaggccaatc aaactcacaa atatatgtca agtttgact 1740

Does Not Comply  
Corrected Diskette Needed

*see p. 5, too*

*insert this mandatory  
numeric identifier  
whenever <221, <222, or  
<223> is shown*

><220>

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/754,853

DATE: 01/23/2001  
TIME: 15:39:47

Input Set : D:\pa\_00330.txt  
Output Set: N:\CRF3\01232001\I754853.raw

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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/754,853

DATE: 01/23/2001  
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RAW SEQUENCE LISTING  
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358	aatttagggat	tattattatt	attattattt	ttattaccat	ccgaagtcgt	tgccgagcga	10020
360	catcgaaaa	ctaccgtgag	aaggaaacag	tagctgttag	cgtgcgcctc	cgactttag	10080
362	cttgcgcgg	agaatgtcg	cggtgagaag	catccggtag	tagcccatgc	tccaccgg	10140
364	cggccagat	ccggtcggag	gccgaaccgg	tttcatttcg	tccggcga	ggtgaacgg	10200
366	gcggcggaa	aacgtgatcg	gaaccggcat	tgcgcgagcc	gttacaaaa	gcaacaaaag	10260
368	cagcaaccgc	caccgcac	agatcgagat	ctggcacttg	cacttattct	gatgcctcg	10320
370	tttaactgtat	ttaagtaacg	attagtgtt	attagtgtt	tgagggtgc	cagtgtgc	10380
372	catcatcgcc	atggatcgta	tcgttgc	cctgtgtgg	tgtgtgtg	tgagagtgg	10440
374	agtgagagtg	agggtggata	aaacaaacaa	acaactatcg	cgcattttgt	tgcgggtgg	10500
376	attagactgt	tactaagtgc	ttaattatgt	gggaaaggaa	agtggatgt	tttagtgg	10560

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/754,853

DATE: 01/23/2001  
TIME: 15:39:48

Input Set : D:\pa\_00330.txt  
Output Set: N:\CRF3\01232001\I754853.raw

378	taacagtaag	tgattattgt	aatatgtat	taggaggaat	aagggtgcaa	cactgcagcg	10620
380	acgaagcgaa	acgtcacgca	cggtggcccc	accatgtctt	tacgtgctt	agaatgaaac	10680
382	ggccctttat	tgccgatgtc	gatttgtctt	tyccactgtg	ggccccccca	catttattat	10740
384	tattccttc	cttttacgaa	ataaaaaata	aaaaatcaaa	caaacaaggc	aaaaggttcc	10800
386	ttaagtattt	agtttcatta	tataaataaa	ataaaatgcct	agatcttagt	aataatcaca	10860
388	ttatgtgggt	tgggtcagga	ataaaagctt	acacacgaaa	aaagaaatct	tgcaagtaaa	10920
390	cagctgaaca	cattaattgt	ttttaaagaa	atctaaagtt	attgaagaaa	acaactgaga	10980
392	catgataatt	tgactaatta	atacttttag	tgaaggagac	gtatttaaa	agataaaagta	11040
394	taattataat	aataattaat	aaaataaaata	acgattaata	tttagtaatt	tcattctatg	11100
396	taatattagt	atgatctcaa	ctcaactgat	aattttcaag	ataatagtt	taattgcact	11160
398	ctgtggaaatc	ttaagttctt	tctccaaaga	aaaaaaaaaa	cattttttct	tcccottgtc	11220
400	gtgttctctt	attctgccat	ctccaattct	gttcacaatc	gtaggttgc	ccgccaatga	11280
402	tgttaatqa	taaagatcaa	atacgtttgc	aatgaatcgg	gatgacaaga	ctgagacaac	11340
404	caatagggtg	agctaaccaa	tgcacaaagt	ctccaaatcaa	taaaacaggc	ccaaaaaggt	11400
406	gggggtgtcc	aaaatgtgaa	ggtaagttt	agttagggtgt	tcacgcctt	gattgcgtct	11460
408	gtgtaaatcc	gtcacccaaat	ccaaacaaaaa	aatattggat	ggattttgtgt	gtttttcttt	11520
410	ttaaatcgac	ctaattctgtat	catgaatgaa	tttgatcgag	atggattttgt	tattaaaaaa	11580
412	agttcaaaaa	taattttctt	aaattttta	aaatattttt	tagaattttac	aatacaatta	11640
414	cttgaatat	agttgcataa	aaaaaattaa	ccaccaattt	caatgcacat	attaactgca	11700
416	tcataaaaatc	aaattgaaaaa	caagtaacca	acaaacattt	aatttataaa	gcaaataata	11760
418	ctaaatcaaa	tttcaaccat	aaagcagata	acaaattgtc	ttgaaaactt	agtaatctta	11820
420	taaagtacac	actagtacaa	aataaactt	aaatcatccc	aaaaaatata	taataactaca	11880
422	atagaaacac	tgcaatata	tgataatgtc	agacaattgc	tcaaccagcc	aacctcacac	11940
424	atagaaacac	ggtaagcaaa	agatcaaaaat	caatttattat	actaataata	aattttaatt	12000
426	atgctatgca	gaaaaaagaa	atatgcwwwaa	aaagaaatca	tatcataaaac	taagttaaaa	12060
428	atattacctt	aagaactaat	agtccatact	cccaatacta	atactcctaa	gaatagtcca	12120
430	agtagtaatc	ctaacactaa	cattatttaa	agtcaaaacca	tacaacttta	aaaaatgttt	12180
432	taaaaaagttc	atcataacat	aatatcaatt	tatattcata	ttgttaacaa	acggaaaaaa	12240
434	aaaaagaaac	tattattgaa	taccttagttc	catcttttt	gttcatctt	attcaactcg	12300
436	taaatcaccg	acattttgct	tattatttt	gagtcaattt	tgggtacaaa	tcaaagcttc	12360
438	aacagtaatg	ggacttaaag	aactacaaaa	atggatcaag	cactcaacct	tttgcactaa	12420
440	atgcagactc	aatgacaca	atagacataa	gaatgacca	tatatctcta	gccatgaaag	12480
442	aaataacatg	atatttggat	gcttcattt	tccaccatgc	caaaatgtca	aatccaaagac	12540
444	cgtcatcttc	attgtcatcc	tttaaataca	tatccaaatc	actcctttgc	tattcaccac	12600
446	atttttatt	cattttcaat	ctaaattgg	cgtcccaatc	ctcatcctca	tcaacatcg	12660
448	tggcattacc	ttgtgaagca	ttgtatgaa	ccaaagtact	agaattacta	ctataatgg	12720
450	aaataggatg	ttctgaagca	tattcaacaa	acattttct	tataagatca	tccaaatttt	12780
452	tcagcatctc	tttggtttgg	tcaacaccat	gcattttctt	aaaacaaaaac	tcaatataat	12840
454	caaatttata	acacagatca	agaaaaagcag	tcacaaataa	aagatagcta	atctgatcac	12900
456	tctctcaata	cttggtaaac	ttgagttgc	tattattgt	ctctttgt	tcacccgatc	12960
458	atccatcat	ctccatotat	ttaggcattt	ctgaatgt	accaacttct	taaagaaatt	13020
460	cttagctgt	acatgtatgt	acccagaaaa	aaaaattgca	tcatagaaaa	cttcaaaaa	13080
462	actcacaaac	acacgagcat	gtttctaaatc	catctcttta	ggacatcctc	cttcactatt	13140
464	tagaagagt	agcacatatg	cagcctcaac	atactcataa	cgattgaaag	tttgtcaaa	13200
466	ttttcagcaa	catctaaat	caaataagt	gagtgacca	aaaaaaa	ttgcatcata	13260
468	aaaaactttc	aaaaaactca	caaacacacq	agcacgttc	taatccatct	ctttaggaca	13320
470	tcctccttca	ctattnagaa	gagtgagcac	atatgcagcc	tcaacatact	cataacgatt	13380
472	gaaagcttgc	tcaaatttcc	agcaacatct	aacatcaa	aagtggagtt	ccatctgg	13440
474	ggcacattaa	gtgttagcat	tgcctttgaa	tttacactaa	cgtcctccgc	acacctctt	13500

**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

FYI

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/754,853

DATE: 01/23/2001  
TIME: 15:39:49

Input Set : D:\pa\_00330.txt  
Output Set: N:\CRF3\01232001\I754853.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No  
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:5859 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 2  
L:6021 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 2  
L:17361 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 3  
L:35814 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#: 4  
L:35816 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#: 4  
L:37292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#: 4  
L:37294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#: 4  
L:37656 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#: 4  
L:37658 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#: 4  
L:37660 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#: 4  
L:37662 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#: 4  
L:37668 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#: 4  
L:44618 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 8  
L:44780 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 8  
L:45076 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 9  
L:45382 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 10  
L:45680 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 11  
L:45985 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 12  
L:46281 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 13  
L:46427 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 14  
L:46589 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 14  
L:46885 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 15  
L:47031 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 16  
L:47193 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 16  
L:47489 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 17  
L:47795 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 18  
L:48095 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 19  
L:48398 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 20  
L:48698 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 21  
L:49001 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 22  
L:49301 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 23